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(FILE 'HOME' ENTERED AT 11:59:50 ON 30 JUL 2002)

FILE 'REGISTRY' ENTERED AT 11:59:57 ON 30 JUL 2002

L1 STRUCTURE UPLOADED
L2 QUE L1
L3 4 SEA SSS FUL L1
D 1-4

FILE 'REGISTRY' ENTERED AT 12:01:48 ON 30 JUL 2002

L4 1 SEA 261775-45-5/RN
D L4 SQIDE TOTAL
D L4 IALL
L5 4748 SEA THIAZ? AND YNYL
D 4748
L6 STRUCTURE UPLOADED
L7 QUE L6
L8 4 SEA SSS FUL L6
D 1-4
L9 1815 SEA THIAZ? YL AND YNYL
D 1815
D 100
L10 1825 SEA (THIAZOLYL OR THIAZOL (2A) YL) AND YNYL
D 1825
L11 STRUCTURE UPLOADED
L12 QUE L11
L13 4 SEA SSS FUL L11
D 1-4
L14 STRUCTURE UPLOADED
L15 QUE L14
L16 2 SEA SSS FUL L14
D 1-2
L17 STRUCTURE UPLOADED
L18 QUE L17
L19 465 SEA SSS FUL L17
D 465
L20 442 SEA L19 NOT BENZOTHIAZOLE
D 442
L21 442 SEA L19 AND L20
L22 23 SEA L19 AND BENZOTHIAZOLE
D 1-23
L23 STRUCTURE UPLOADED
L24 QUE L23
L25 116 SEA SSS FUL L23
D 116
L26 STRUCTURE UPLOADED
L27 QUE L26
L28 22 SEA SSS FUL L26
D 1-22
L29 578 SEA L8 OR L13 OR L16 OR L20 OR L25 OR L28
L30 11 SEA L29 AND SALT
D 1-11

FILE 'CAPLUS, EMBASE, BIOSIS, DRUGU, CAOLD, USPATFULL, IFIPAT' ENTERED
AT

12:26:28 ON 30 JUL 2002

L31 10 SEA L30
D 1-10
D 1-10 HITSTR
D 10 IALL
D 10
D 10 HITSTR
D 7 IALL HITSTR

L32 7 SEA L30 AND (SALT OR SALTS)
D 7 KWIC
D 7

L33 68 SEA L29 NOT L30

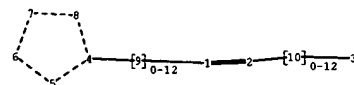
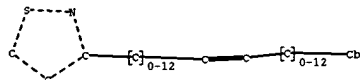
L34 20 SEA L33 AND (SALT OR SALTS)
D 1-20 KWIC
D 1-20
D 1-20 HITSTR
D 1,2,3,6,15,16,17,18
D 6 IALL HITSTR
D 1-3

L35 6 SEA L33 AND (RECEPTOR? OR GLUTAMATE OR METABOTROPIC OR
MODULAT?)
D 1-6
D 6 KWIC
D 6 HITSTR
D 5 HITSTR
D 6
D 6 HITSTR
D L34 15,16,17,18
D L34 18
D L34 18 HITSTR
D L34 117
D L34 17 HITSTR
D L34 16
D L34 16 HITSTR
D L34 15
D L34 15 HITSTR

L36 48 SEA L33 NOT L34

L37 46 SEA L36 NOT L35
D 1-46
D 46 IALL
D 45 IALL
D 1-44 HITSTR
D 1-10
D 1,6,8,10,12,16,20,22,32,33,36,37,38,39,40,41,42,43,44 IALL

HI



chain nodes :

1 2 3 9 10

ring nodes :

4 5 6 7 8

chain bonds :

1-2 1-9 2-10 3-10 4-9

ring bonds :

4-5 4-8 5-6 6-7 7-8

exact/norm bonds :

4-5 4-8 5-6 6-7 7-8

exact bonds :

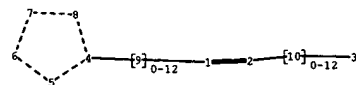
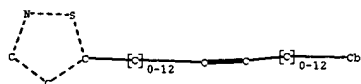
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Match level :

1:CLASS 2:CLASS 3:Atom 4:Atom 5:Atom 6:Atom 7:Atom 8:Atom 9:CLASS
10:CLASS

Generic attributes :

3:
Saturation : Unsaturated



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ring nodes :

4 5 6 7 8

chain bonds :

1-2 1-9 2-10 3-10 4-9

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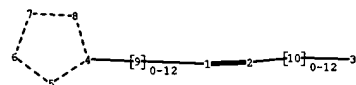
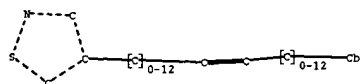
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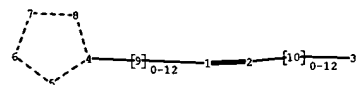
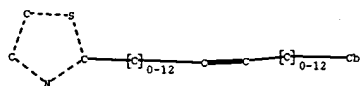
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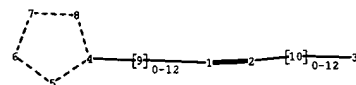
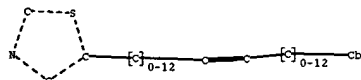
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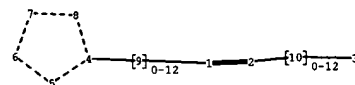
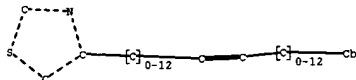
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